

Art Unit: 1652

(i) In claim 15 line 2, please delete the word "therapeutic" and in line 8 delete the phrase "the following"

(ii) In claims 15 and 27 part (b), lines ~~2 and 4~~, replace "high-stringent condition with a complement" with "high-stringency conditions with the full complement".

*In claims 15 and 27 part (b) line 4 replace "high-stringent condition" with "high-stringency conditions with the full complement"*

(iii) In claim 27, line 5, delete "the following"

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(iv) Please amend claim 28 as follows:

28. The diagnostic method according to claim 27, wherein the expression level is assayed using a probe that is a nucleic acid capable of hybridizing with a nucleic acid having the nucleotide sequence shown in SEQ ID NO: 8 under high-stringen[t]cy conditions or [a] the full complement thereof [as a probe or primer], wherein hybridization under high-stringen[t]cy conditions is carried out by hybridizing at 65° C for 16 hours in a hybridization solution (6xSSC, 0.5% SDS, 5xDenhardt's solution, 100µg/ml salmon sperm DNA), washing at 65 °C for 5 minutes in a washing solution (6xSSC, 0.5%SDS), and thereafter washing twice at 65° C for 30 min in washing solution (0.1xSSC).